☐1: O14734. Peroxisomal acyl-...[gi:14285685] Links

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LOCUS       PTE1_HUMAN              319 aa            linear   PRI 15-JUN-2002
DEFINITION  Peroxisomal acyl-coenzyme A thioester hydrolase 1 (Peroxisomal
            long-chain acyl-coA thioesterase 1) (HIV-Nef associated acyl coA
            thioesterase) (Thioesterase II) (hTE).
ACCESSION   O14734
VERSION     O14734  GI:14285685
DBSOURCE    swissprot: locus PTE1_HUMAN, accession O14734;
            class: standard.
            extra accessions:O15261,created: Oct 16, 2001.
            sequence updated: Oct 16, 2001.
            annotation updated: Jun 15, 2002.
            xrefs: gi: 2318124, gi: 2318125, gi: 2243145, gi: 2243146, gi:
            4680312, gi: 4680313, gi: 13929448, gi: 3191970
            xrefs (non-sequence databases): HSSP P23911, InterPro IPR003703,
            Pfam PF02551
KEYWORDS    Hydrolase; Serine esterase; Peroxisome.
SOURCE      Homo sapiens.
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (residues 1 to 319)
  AUTHORS   Watanabe,H., Shiratori,T., Shoji,H., Miyatake,S., Okazaki,Y.,
            Ikuta,K., Sato,T. and Saito,T.
  TITLE     A novel acyl-CoA thioesterase enhances its enzymatic activity by
            direct binding with HIV Nef
  JOURNAL   Biochem. Biophys. Res. Commun. 238 (1), 234-239 (1997)
  MEDLINE   97445158
  REMARK    SEQUENCE FROM N.A.
REFERENCE   2  (residues 1 to 319)
  AUTHORS   Liu,L.X., Margottin,F., Le Gall,S., Schwartz,O., Selig,L.,
            Benarous,R. and Benichou,S.
  TITLE     Binding of HIV-1 Nef to a novel thioesterase enzyme correlates with
            Nef-mediated CD4 down-regulation
  JOURNAL   J. Biol. Chem. 272 (21), 13779-13785 (1997)
  MEDLINE   97298085
  REMARK    SEQUENCE FROM N.A.
            TISSUE=Lymphoid
REFERENCE   3  (residues 1 to 319)
  AUTHORS   Jones,J.M., Nau,K., Geraghty,M.T., Erdmann,R. and Gould,S.J.
  TITLE     Identification of peroxisomal acyl-CoA thioesterases in yeast and
            humans
  JOURNAL   J. Biol. Chem. 274 (14), 9216-9223 (1999)
  MEDLINE   99194760
  REMARK    SEQUENCE FROM N.A.
            TISSUE=Muscle
REFERENCE   4  (residues 1 to 319)
  AUTHORS   Deloukas,P., Matthews,L.H., Ashurst,J., Burton,J., Gilbert,J.G.R.,
            Jones,M., Stavrides,G., Almeida,J.P., Babbage,A.K., Bagguley,C.L.,
            Bailey,J., Barlow,K.F., Bates,K.N., Beard,L.M., Beare,D.M.,
            Beasley,O.P., Bird,C.P., Blakey,S.E., Bridgeman,A.M., Brown,A.J.,
            Buck,D., Burrill,W.D., Butler,A.P., Carder,C., Carter,N.P.,
            Chapman,J.C., Clamp,M., Clark,G., Clark,L.N., Clark,S.Y.,
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Clee,C.M., Clegg,S., Cobley,V.E., Collier,R.E., Connor,R.E., Corby,N.R., Coulson,A., Coville,G.J., Deadman,R., Dhami,P.D., Dunn,M., Ellington,A.G., Frankland,J.A., Fraser,A., French,L., Garner,P., Grafham,D.V., Griffiths,C., Griffiths,M.N.D., Gwilliam,R., Hall,R.E., Hammond,S., Harley,J.L., Heath,P.D., Ho,S., Holden,J.L., Howden,P.J., Huckle,E., Hunt,A.R., Hunt,S.E., Jekosch,K., Johnson,C.M., Johnson,D., Kay,M.P., Kimberley,A.M., King,A., Knights,A., Laird,G.K., Lawlor,S., Lehvaslaiho,M.H., Leversha,M.A., Lloyd,C., Lloyd,D.M., Lovell,J.D., Marsh,V.L., Martin,S.L., McConnachie,L.J., McLay,K., McMurray,A.A., Milne,S.A., Mistry,D., Moore,M.J.F., Mullikin,J.C., Nickerson,T., Oliver,K., Parker,A., Patel,R., Pearce,T.A.V., Peck,A.I., Phillimore,B.J.C.T., Prathalingam,S.R., Plumb,R.W., Ramsay,H., Rice,C.M., Ross,M.T., Scott,C.E., Sehra,H.K., Shownkeen,R., Sims,S., Skuce,C.D., Smith,M.L., Soderlund,C., Steward,C.A., Sulston,J.E., Swann,R.M., Sycamore,N., Taylor,R., Tee,L., Thomas,D.W., Thorpe,A., Tracey,A., Tromans,A.C., Vaudin,M., Wall,M., Wallis,J.M., Whitehead,S.L., Whittaker,P., Willey,D.L., Williams,L., Williams,S.A., Wilming,L., Wray,P.W., Hubbard,T., Durbin,R.M., Bentley,D.R., Beck,S. and Rogers,J.

TITLE The DNA sequence and comparative analysis of human chromosome 20
JOURNAL Nature 414 (6866), 865-871 (2001)
MEDLINE 21638749
REMARK SEQUENCE FROM N.A.

COMMENT ----------

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[FUNCTION] MAY PLAY A ROLE IN FATTY ACID OXIDATION RATHER THAN FORMATION OF FATTY ACIDS. MAY MEDIATE NEF-INDUCED DOWN-REGULATION OF CD4.
[CATALYTIC ACTIVITY] Palmitoyl-CoA + H(2)O = CoA + palmitate.
[SUBUNIT] INTERACTS WITH HIV-1 NEF.
[SUBCELLULAR LOCATION] Peroxisomal.
[SIMILARITY] BELONGS TO THE C/M/P THIOESTER HYDROLASE FAMILY.

FEATURES Location/Qualifiers

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/db_xref="taxon:9606"

gene 1..319
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Protein 1..319
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Site 232
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Region 291..293
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Site 317..319
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                     /gene="PTE1"
                     /region_name="Conflict"
                     /note="L -> R (IN REF. 2)."
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      121 ificqasfqq aqpspmqhqf smptvpppee lldcetlidq ylrdpnlqkr yplalnriaa
      181 qevpieikpv npsplsqlqr mepkqmfwvr argyigegdm kmhccvaayi sdyaflgtal
      241 lphqwqhkvh fmvsldhsmw fhapfradhw mlyecespwa ggsrglvhgr lwrqdgvlav
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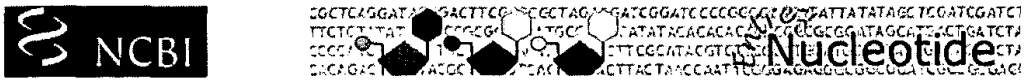
Revised: July 5, 2002.



☐1: R35332. yg65c03.r1 Soares...[gi:792233] Links

IDENTIFIERS

dbEST Id: **205393**
EST name: yg65c03.r1
GenBank Acc: R35332
GenBank gi: 792233
GDB Id: 410808

CLONE INFO
Clone Id: IMAGE:38267 (5')
Insert length: 1325
DNA type: cDNA

PRIMERS
Sequencing: M13RP1
PolyA Tail: Unknown

SEQUENCE

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CGGCGATCCCCCTGGGACCTCCGTAGCGTCTTGGTCACGACCGTGCTCAACCTCGAGCCG
CTGGACGAGGATCTCTTCAGAGGAAGGCATTACTGGGTACCGGCCAAGAGGCTGTTTGGT
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TGCACTCCCTGCACTGCTACTTTGTTCGGGCAGGGGACCCGAAGCTGCCAGTACTGTACC
AAGTGGAGCGGACACGAACAGGGTCGAGCTTCTTCGGTGCGCTCTGTGAAGGCCGTGCAA
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ATTGCAGCACCAGT

Quality: High quality sequence stops at base: 327

Entry Created: May 2 1995
Last Updated: May 2 1995

COMMENTS

Insert Size: 1325
High quality sequence stops: 327 Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL ; contact the IMAGE Consortium (info@image.llnl.gov) for further information.

PUTATIVE ID Assigned by submitter
SP:TESB_ECOLI P23911 ACYL-COA THIOESTERASE II ;

LIBRARY
Lib Name: Soares infant brain 1NIB
Organism: Homo sapiens
Sex: female
Organ: whole brain
Develop. stage: 73 days post natal
Lab host: DH10B (ampicillin resistant)
Vector: Lafmid BA
R. Site 1: Not I
R. Site 2: Hind III
Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

[5' AACTGGAAGAATTCGCGGCCGCAGGAATTTTTTTTTTTTTTTT 3'];
double-stranded cDNA was ligated to Hind III adaptors
(Pharmacia), digested with Not I and directionally cloned
into the Not I and Hind III sites of the Lafmid BA vector.
Library went through one round of normalization. Library
constructed by Bento Soares and M.Fatima Bonaldo.

SUBMITTER

Name: Wilson RK
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E-mail: est@watson.wustl.edu

CITATIONS

Title: The WashU-Merck EST Project
Authors: Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M. , Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P., Wilson,R.
Year: 1995
Status: Unpublished

MAP DATA

Revised: July 5, 2002.

WEST

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Search Results -

Terms	Documents
L3 and @RLAD<19980101	100

Database: US Patents Full-Text Database; US Pre-Grant Publication Full-Text Database; JPO Abstracts Database; EPO Abstracts Database; Derwent World Patents Index; IBM Technical Disclosure Bulletins

Search: L4

Refine Search | Recall Text | Clear

Search History

DATE: Sunday, October 06, 2002 Printable Copy Create Case

Set Name side by side	Query	Hit Count	Set Name result set
DB=USPT; THES=ASSIGNEE; PLUR=YES; OP=OR			
L4	L3 and @RLAD<19980101	100	L4
L3	l1 and antibody	151	L3
L2	L1 and peroxisom$	23	L2
L1	thioesterase	251	L1

END OF SEARCH HISTORY

FILE 'MEDLINE, CAPLUS, EMBASE, BIOSIS' ENTERED AT 14:40:29 ON 06 OCT 2002

L1	347	S EC 3.1.2.2
L2	281	DUP REM L1 (66 DUPLICATES REMOVED)
L3	2313	S THIOESTERASE OR (THIOESTER (1W) HYDROLASE)
L4	11	S L3 AND PTE1
L5	3	DUP REM L4 (8 DUPLICATES REMOVED)
L6	154	S L3 AND PEROXISOM?
L7	59	DUP REM L6 (95 DUPLICATES REMOVED)
L8	9	S L7 AND PY<1998